

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/798,025
Source: IFWO
Date Processed by STIC: 12/14/05

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 12/14/2005

PATENT APPLICATION: US/10/798,025

TIME: 18:25:29

Input Set : N:\Cr4\12132005\K798025.raw

Output Set: N:\CRF4\12142005\J798025.raw

1 <110> APPLICANT: Langmore, John
 2 Makarov, Vladimir
 3 <120> TITLE OF INVENTION: Method of Producing a DNA Library Using Positional
 Amplification
 4 <130> FILE REFERENCE: UMIC:047US0/10103482
 C--> 5 <140> CURRENT APPLICATION NUMBER: US/10/798,025
 6 <141> CURRENT FILING DATE: 2004-03-11
 7 <160> NUMBER OF SEQ ID NOS: 121
 8 <170> SOFTWARE: PatentIn version 3.1
 10 <210> SEQ ID NO: 1
 11 <211> LENGTH: 24
 12 <212> TYPE: DNA
 13 <213> ORGANISM: Unknown
 14 <220> FEATURE:
 15 <223> OTHER INFORMATION: Primer
 16 <400> SEQUENCE: 1
 17 gatcgcttat acctaggacc atgt 24
 19 <210> SEQ ID NO: 2
 20 <211> LENGTH: 22
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Artificial Sequence
 23 <220> FEATURE:
 24 <223> OTHER INFORMATION: DNA/RNA Primer
 25 <220> FEATURE:
 26 <221> NAME/KEY: misc_feature
 27 <222> LOCATION: (1)..(22)
 28 <223> OTHER INFORMATION: Ribonucleotide at 7, 10 ,13, 19
 29 <400> SEQUENCE: 2
 30 gttacauggu ccuaggtaua gg 22
 32 <210> SEQ ID NO: 3
 33 <211> LENGTH: 23
 34 <212> TYPE: DNA
 35 <213> ORGANISM: Unknown
 36 <220> FEATURE:
 37 <223> OTHER INFORMATION: Primer
 38 <400> SEQUENCE: 3
 39 gttacatggt cctaggtata ggc 23
 41 <210> SEQ ID NO: 4
 42 <211> LENGTH: 37
 43 <212> TYPE: DNA
 44 <213> ORGANISM: Unknown
 45 <220> FEATURE:
 46 <223> OTHER INFORMATION: Primer
 47 <400> SEQUENCE: 4

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51 <211> LENGTH: 45
52 <212> TYPE: DNA
53 <213> ORGANISM: Unknown
54 <220> FEATURE:
55 <223> OTHER INFORMATION: DNA/RNA Primer
56 <220> FEATURE:
57 <221> NAME/KEY: misc_feature
58 <222> LOCATION: (1)..(45)
59 <223> OTHER INFORMATION: Ribonucleotide at 24, 29, 35, 41
60 <400> SEQUENCE: 5
61      aggtcgccgc cctgatgaat tcgutacaug gtccuaggta uaggc      45
63 <210> SEQ ID NO: 6
64 <211> LENGTH: 12
65 <212> TYPE: DNA
66 <213> ORGANISM: Unknown
67 <220> FEATURE:
68 <223> OTHER INFORMATION: Primer
69 <400> SEQUENCE: 6
70      gggcgggcgac ct      12
72 <210> SEQ ID NO: 7
73 <211> LENGTH: 25
74 <212> TYPE: DNA
75 <213> ORGANISM: Unknown
76 <220> FEATURE:
77 <223> OTHER INFORMATION: Primer
78 <400> SEQUENCE: 7
79      gggagatctg aattcccccc ccccc      25
81 <210> SEQ ID NO: 8
82 <211> LENGTH: 23
83 <212> TYPE: DNA
84 <213> ORGANISM: Unknown
85 <220> FEATURE:
86 <223> OTHER INFORMATION: Primer
87 <400> SEQUENCE: 8
88      gggagatctg aattcaaaaa aaa      23
90 <210> SEQ ID NO: 9
91 <211> LENGTH: 24
92 <212> TYPE: DNA
93 <213> ORGANISM: Unknown
94 <220> FEATURE:
95 <223> OTHER INFORMATION: Primer
96 <400> SEQUENCE: 9
97      gaattcagat ctcccggggtc accg      24
99 <210> SEQ ID NO: 10
100 <211> LENGTH: 30
101 <212> TYPE: DNA
102 <213> ORGANISM: Unknown

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103 <220> FEATURE:
104 <223> OTHER INFORMATION: Primer
105 <400> SEQUENCE: 10
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108 <210> SEQ ID NO: 11
109 <211> LENGTH: 30
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111 <213> ORGANISM: Unknown
112 <220> FEATURE:
113 <223> OTHER INFORMATION: Primer
114 <400> SEQUENCE: 11
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118 <211> LENGTH: 42
119 <212> TYPE: DNA
120 <213> ORGANISM: Unknown
121 <220> FEATURE:
122 <223> OTHER INFORMATION: Primer
123 <400> SEQUENCE: 12
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126 <210> SEQ ID NO: 13
127 <211> LENGTH: 25
128 <212> TYPE: DNA
129 <213> ORGANISM: Unknown
130 <220> FEATURE:
131 <223> OTHER INFORMATION: Primer
132 <400> SEQUENCE: 13
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135 <210> SEQ ID NO: 14
136 <211> LENGTH: 25
137 <212> TYPE: DNA
138 <213> ORGANISM: Unknown
139 <220> FEATURE:
140 <223> OTHER INFORMATION: Primer
141 <400> SEQUENCE: 14
142      gcggtgaccc gggagatctg aattc                   25
144 <210> SEQ ID NO: 15
145 <211> LENGTH: 38
146 <212> TYPE: DNA
147 <213> ORGANISM: Unknown
148 <220> FEATURE:
149 <223> OTHER INFORMATION: Primer
150 <400> SEQUENCE: 15
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153 <210> SEQ ID NO: 16
154 <211> LENGTH: 27
155 <212> TYPE: DNA
156 <213> ORGANISM: Unknown
157 <220> FEATURE:

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158 <223> OTHER INFORMATION: Primer
159 <220> FEATURE:
160 <221> NAME/KEY: misc_feature
161 <222> LOCATION: (27)..(27)
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167 <211> LENGTH: 23
168 <212> TYPE: DNA
169 <213> ORGANISM: Artificial Sequence
170 <220> FEATURE:
171 <223> OTHER INFORMATION: DNA/RNA Primer
172 <220> FEATURE:
173 <221> NAME/KEY: misc_feature
174 <222> LOCATION: (1)..(23)
175 <223> OTHER INFORMATION: Ribonucleotide at 7, 10, 13, 19
176 <220> FEATURE:
177 <221> NAME/KEY: misc_feature
178 <222> LOCATION: (23)..(23)
179 <223> OTHER INFORMATION: n equals ddC
180 <400> SEQUENCE: 17
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184 <211> LENGTH: 26
185 <212> TYPE: DNA
186 <213> ORGANISM: Unknown
187 <220> FEATURE:
188 <223> OTHER INFORMATION: Primer
189 <400> SEQUENCE: 18
190      gatcgccctat acctaggacc atgtaa      26
192 <210> SEQ ID NO: 19
193 <211> LENGTH: 23
194 <212> TYPE: DNA
195 <213> ORGANISM: Artificial Sequence
196 <220> FEATURE:
197 <223> OTHER INFORMATION: DNA/RNA Primer
198 <220> FEATURE:
199 <221> NAME/KEY: misc_feature
200 <222> LOCATION: (1)..(23)
201 <223> OTHER INFORMATION: Ribonucleotide at 7, 10, 13, 19
202 <400> SEQUENCE: 19
203      gttacauggu ccuaggtaua ggc      23
205 <210> SEQ ID NO: 20
206 <211> LENGTH: 37
207 <212> TYPE: DNA
208 <213> ORGANISM: Unknown
209 <220> FEATURE:
210 <223> OTHER INFORMATION: Primer

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Input Set : N:\Crf4\12132005\K798025.raw
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211 <400> SEQUENCE: 20
212      gatcgccctat acctaggacc atgtaacgaa ttcatca
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216 <212> TYPE: DNA
217 <213> ORGANISM: Artificial Sequence
218 <220> FEATURE:
219 <223> OTHER INFORMATION: DNA/RNA Primer
220 <220> FEATURE:
221 <221> NAME/KEY: misc_feature
222 <222> LOCATION: (1)..(45)
223 <223> OTHER INFORMATION: Ribonucleotide at 24, 29, 35, 41
224 <400> SEQUENCE: 21
225      aggtcgccgc cctgatgaat tcgutacaug gtccuaggta uaggc
227 <210> SEQ ID NO: 22
228 <211> LENGTH: 26
229 <212> TYPE: DNA
230 <213> ORGANISM: Unknown
231 <220> FEATURE:
232 <223> OTHER INFORMATION: Primer
233 <220> FEATURE:
234 <221> NAME/KEY: misc_feature
235 <222> LOCATION: (26)..(26)
236 <223> OTHER INFORMATION: n equals ddC
237 <400> SEQUENCE: 22
W--> 238      gggagatctg aattcccccc ccccn
240 <210> SEQ ID NO: 23
241 <211> LENGTH: 25
242 <212> TYPE: DNA
243 <213> ORGANISM: Unknown
244 <220> FEATURE:
245 <223> OTHER INFORMATION: Primer
246 <220> FEATURE:
247 <221> NAME/KEY: misc_feature
248 <222> LOCATION: (25)..(25)
249 <223> OTHER INFORMATION: n equals ddC
250 <400> SEQUENCE: 23
W--> 251      gaattcagat ctcccggtc accgn
253 <210> SEQ ID NO: 24
254 <211> LENGTH: 53
255 <212> TYPE: DNA
256 <213> ORGANISM: Unknown
257 <220> FEATURE:
258 <223> OTHER INFORMATION: Primer
259 <400> SEQUENCE: 24
260      gttacatggt cctaggtata ggcgcggtga cccgggagat ctgccccccc ccc
262 <210> SEQ ID NO: 25
263 <211> LENGTH: 42
264 <212> TYPE: DNA

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RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : N:\Cr4\12132005\K798025.raw
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:16; N Pos. 27
Seq#:17; N Pos. 23
Seq#:22; N Pos. 26
Seq#:23; N Pos. 25
Seq#:26; N Pos. 25
Seq#:27; N Pos. 25
Seq#:42; N Pos. 47,48,49,50
Seq#:44; N Pos. 47,48,49,50
Seq#:62; N Pos. 47,48,49,50
Seq#:64; N Pos. 26

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 3

VERIFICATION SUMMARY

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Input Set : N:\Crf4\12132005\K798025.raw

Output Set: N:\CRF4\12142005\J798025.raw

L:5 M:270 C: Current Application Number differs, Replaced Current Application Number

L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0

L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0

L:238 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0

L:251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0

L:282 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0

L:299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0

L:447 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:0

L:469 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:0

L:644 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62 after pos.:0

L:666 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:0